

# Cyanogen Bromide (CNBr) Cleavage

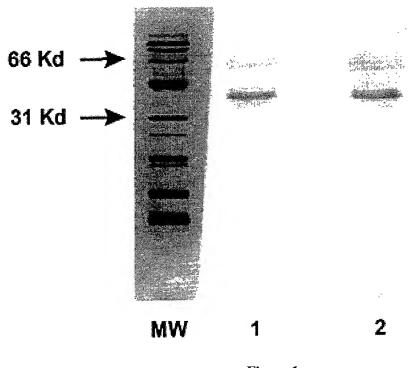
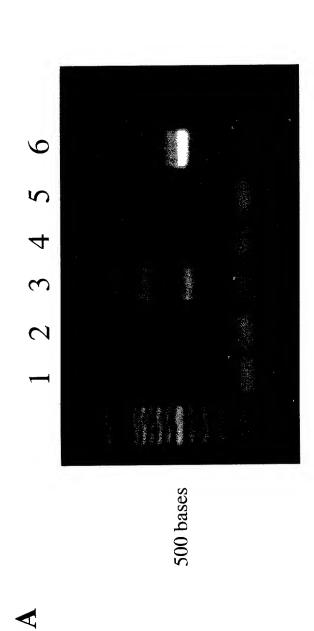


Figure 1



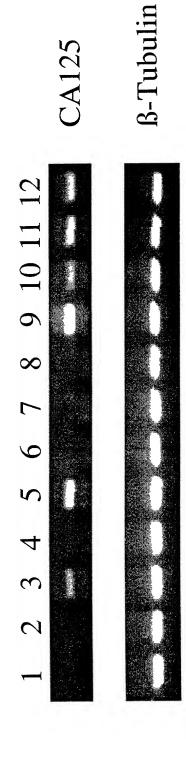
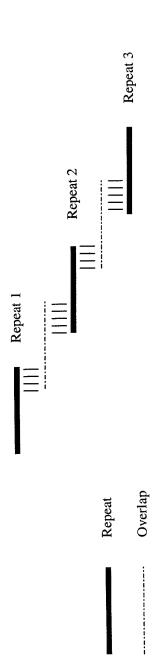


Figure 2

# A Strategy for Placing Repeat Sequences in Contiguous Order Using Overlap Sequence Allignment



156 314 470 625 781 1 ATOPENVIPTITINLOYBEDMRHPGSRKFNATERELGGLIKPLFRUSSLEYLYSGCRLASIAPEKDSSAMAVDAICTHRPDPEDLGLDRERLYWELSNLTNGIQELGFYTLDRNSLYVNGFTHRSMPTTSTPGTSTVDVGTSGFPSSSPSPT

1 ATOPENAPETINFTITINLOYBEDMRTGSRKFNTTMESVLQGLIAFLERNSTSERLYBELTERPEKDGAATGYDAICTHRLDPRSPGINREQLYWELSKLTNDIEELGPYTLDRNSLYVNGFTHGSSVSTTSTPGTSTVDLGTSSFPTIM

1157 AAGPLIAMPFTINLOYGEDMGHPGSRKFNTTTBTVLGGLIAPIFKNTSVGLISSERDGAATGVDAICTHRLDPRSPGINRERLYWELSQLTNGIKELGPYTLDRNSLYVNGFTHWIPVPTSSTPGTSTVDLGTSGFPSLESPT

471 TAGPLIVLFTINFITTINLEXPEDMHRPGSRKFNTTERVLGTLLGPMFKNTSVGLLYSGCRLTLIRSEKDGAATGVDAICTHRLDPRSPGUNWELSQLTNGIKELGPYTLDRNSLYVNGFTHWIPVPTSSTPGTSTVDLGTSGFPSSLESPT

626 AAGPLIVPFTLNFTITINLQYBEDMHHPGSRKFNTTERVLGGLLGPMFKNTSVGLLYSGCRLTLIRSEKDGAATGVDAICTHRLDPRSRPQVPREQLTNGIKELGPYTLDRNSLYVNGFTHQTSAPNTSTPGTSTVDLGTSGFPSSLESPT

626 AAGPLIVPFTLNFTITINLQYBEDMHHPGSRKFNTTERVLGGLLGPMFKNTSVGLLYSGCRLTLIRSEKDGAATGVDAICTHRLDPRSRPQVPREQLTNGIKELGPYTLDRNSLYVNGFTHQTSAPNTSTPGTSTVDLGTSGFPSSLPSPT (SEQ ID NO: 159)

Figure 3 (SEQ ID NOS: 158, 159, 160, and 161)

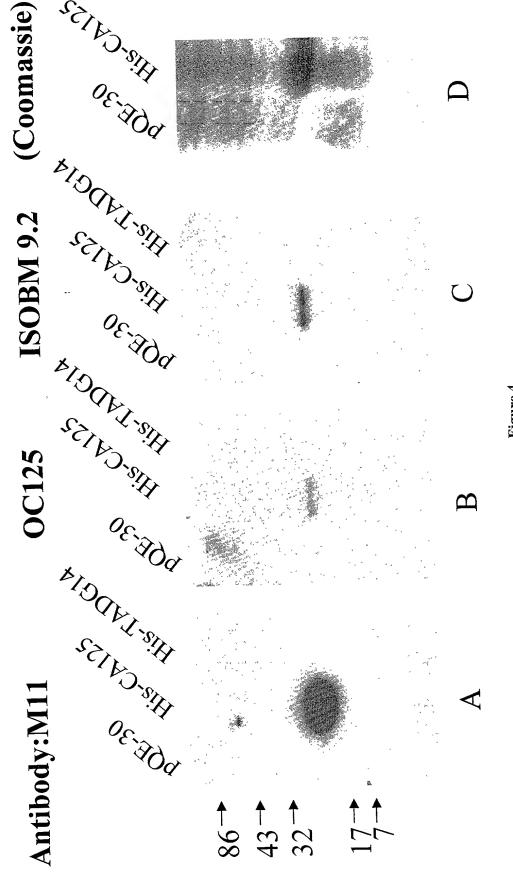
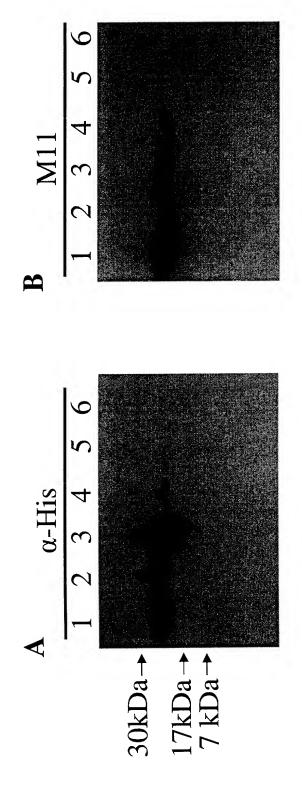
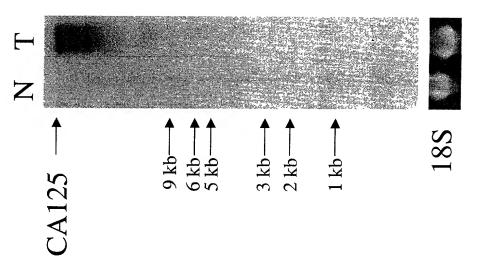


Figure 4



0 0 00 QLTNSITELG PYTLDRDSLY VNGFNPRSSV PTTSTPGTST VHLATSGTPS C- SVGPLYSGCR LTLLRPEKHE AATGVDTICT HRVDPIGPGL DRERLYWELS EPGPLLIPFT FNFTITNLHY EENMQHPGSR KFNTTERVLQ GLLKPLFKNT 0 0000 00 SLPKLT 51 101 151

Figure 5 (SEQ ID NO: 150)



. A. .

Figure 6

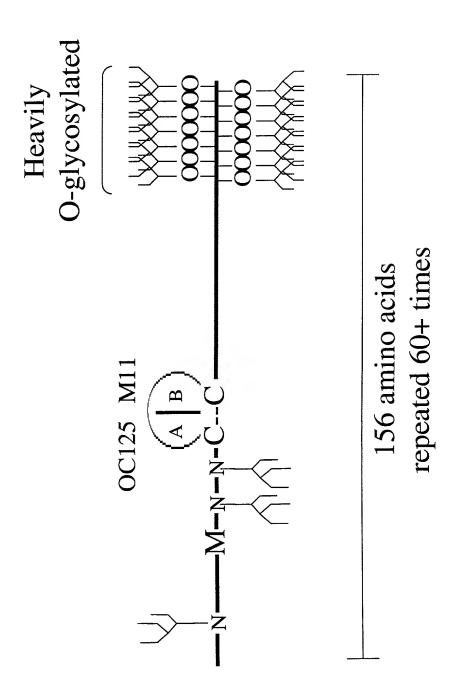
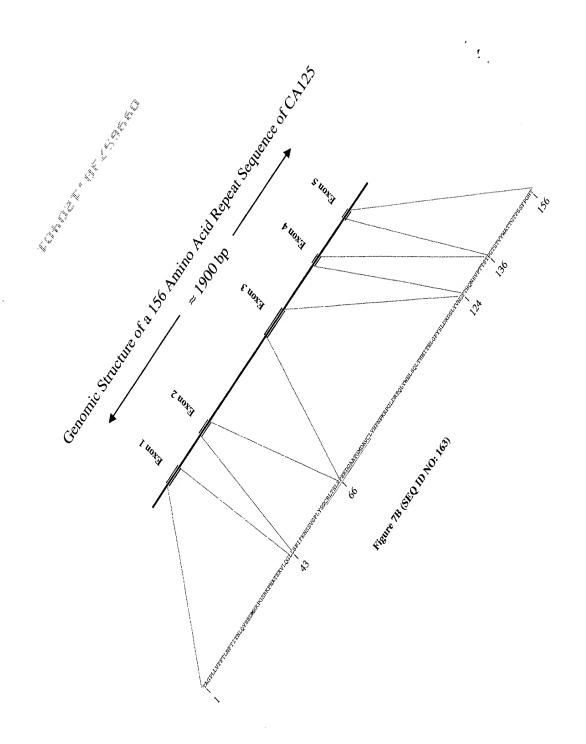


Figure 7A



### Exon 1

ATVPFMVPFTLNFTITNLQYEEDMRHPGSRKFNATERELQGL (SEQ ID NO: 164) TAVPLLVPFTLNFTITNLQYGEDMRHPGSRKFNTTERVLQGL (SEQ ID NO: 165) VPGPLLVPFTLNFTITNLQYEEAMRHPGSRKFNTTERVLQGL (SEQ ID NO: 166) APGPLLVPFTLNFTITNLQYEEDMRHPGSRKFSTTERVLQGL (SEQ ID NO: 167) APGPLLVPFTLNFTITNLQYEEDMRHPGSRKFNTTERVLQGL (SEQ ID NO: 168) APGPLLVPFTLNFTITNLQYEVDMRHPGSRKFNTTERVLQGL (SEQ ID NO: 169) SAGPLLVPFTLNFTITNLQYEEDMRHPGSRKFNTTERVLQGL (SEQ ID NO: 170) AAGPLLMPFTLNFTITNLQYEEDMRRTGSRKFNTMESVLQGL (SEQ ID NO: 171) TASPLLVLFTINCTITNLQYEEDMRRTGSRKFNTMESVLQGL (SEQ ID NO: 172) AAGPLLVPFTLNFTITNLQYGEDMGHPGSRKFNTTERVLQGL (SEQ ID NO: 173) TAGPLLIPFTLNFTITNLQYGEDMGHPGSRKFNTTERVLQGL (SEQ ID NO: 174) TAGPLLVPFTLNFTITNLQYGEDMGHPGSRKFNTTERVLQGL (SEQ ID NO: 175) TAGPLLVLFTLNFTITNLKYEEDMHRPGSRKFNTTERVLQTL (SEQ ID NO: 176) TAGPLLVPFTLNFTITNLQYEEDMHRPGSRKFNATERVLQGL (SEQ ID NO: 177) TAGPLLVPFTLNFTITNLQYEEDMHRPGSRRFNTTERVLQGL (SEQ ID NO: 178) TAGPLLVPFTLNFTITNLQYEEDMHRPGSRKFNTTERVLQGL (SEQ ID NO: 179) APVPLLIPFTLNFTITNLQYEEDMHRPGSRKFNTTERVLQGL (SEQ ID NO: 180) ATGPVLLPFTLNFTITNLQYEEDMHRPGSRKFNTTERVLQGL (SEQ ID NO: 181) AAGPLLVPFTLNFTITNLQYEEDMHHPGSRKFNTTERVLQGL (SEQ ID NO: 182) SAGPLLVPFTLNFTITNLQYEEDMHHPGSRKFNTTERVLQGL (SEQ ID NO: 183) TASPLLVLFTINFTITNQRYEENMHHPGSRKFNTTERVLQGL (SEQ ID NO: 184) TASPLLVLFTINFTITNLRYEENMHPGSRKFNTTERVLQGL (SEQ ID NO: 185) EPGPLLIPFTFNFTITNLHYEENMQHPGSRKFNTTERVLQGL (SEQ ID NO: 186) EPGPLLIPFTFNFTITNLRYEENMQHPGSRKFNTTERVLQGL (SEQ ID NO: 187) APVPLLIPFTLNFTITNLHYEENMQHPGSRKFNTTERVLQGL (SEQ ID NO: 188) APVPLLIPFTLNFTITDLHYEENMQHPGSRKFNTTERVLQGL (SEQ ID NO: 189) AASPLLVLFTLNGTITNLRYEENMQHPGSRKFNTTERVLQGL (SEQ ID NO: 190) TAGPLLVPFTLNFTITNLKYEEDMHCPGSRKFNTTERVLQSL (SEQ ID NO: 191) AASHLLILFTLNFTITNLRYEENMW.PGSRKFNTTERVLQGL (SEQ ID NO: 192) TGVVSEEPFTLNFTINNLRYMADMGQPGSLKFNITDNVMKHL (SEQ ID NO: 193) AMGYHLKTLTLNFTISNLQYSPDMGKGSATFNSTEGVLQHLL (SEQ ID NO: 194)

### Exon 2

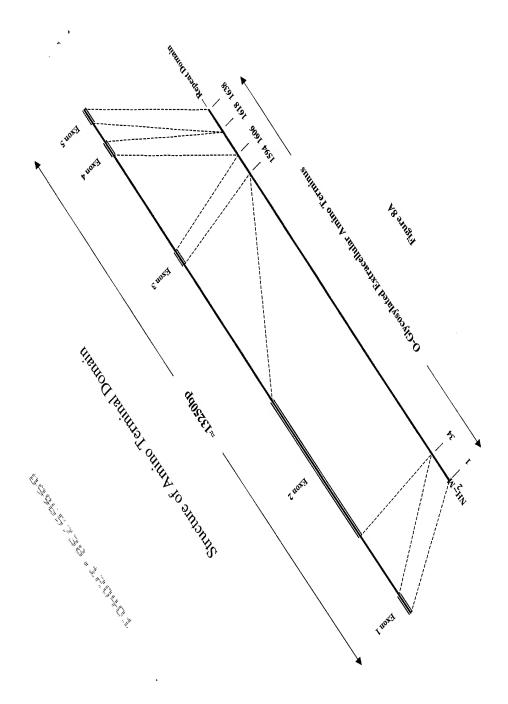
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LKPLFRNSSLEYLYSGCRLASLR	(SEQ	ID	NO:	195
LKPLFKNTSVSSLYSGCRLTLLR	(SEQ	ID	NO:	196
LKPLFKNTSVGPLYSGCRLTLLR	(SEQ	ID	NO:	197
LKPLFKSTSVGPLYSGCRLTLLR	(SEQ	ID	NO:	198
LKPLFKSTSVGPLYSSCRLTLLR	(SEQ	ID	NO:	199
LKPLFKNTSVGPLYSGCRLTSLR	(SEQ	ID	NO:	200
LGPIFKNTSVGPLYSGCRLTSLR	(SEQ	ID	NO:	201
LGPMFKNTSVGLLYSGCRLTLLR	(SEQ	ID	NO:	202
LGPMFKNTSVGPLYSGCRLTLLR	(SEQ	ID	NO:	203
LGPMFKNTSVGPLYSGCRLTSLR	(SEQ	ID	NO:	204)
LGPLFKNSSVGPLYSGCRLISLR	(SEQ	ID	NO:	205)
LGPLFKNSSVDPLYSGCRLTSLR	(SEQ	ID	NO:	206)
LSPIFKNSSVGPLYSGCRLTSLR	(SEQ	ID	NO:	207)
LSPIFKNTSVGPLYSGCRLTLLR	(SEQ	ID	NO:	208)
LSPLFQRSSLGARYTGCRVIALR	(SEQ	ID	NO:	209)
LRPLFKNTSVSSLYSGCRLTLLR	(SEQ	ID	NO:	210)
LRPLFKNTSVGPLYSGSRLTLLR	(SEQ	ID	NO:	211)
LRPLFKNTSIGPLYSSCRLTLLR	(SEQ	ID	NO:	212)
LRPLFKSTSVGPLYSGCRLTLLR	(SEQ	ID	NO:	213)
LRPVFKNTSVGLLYSGCRLTLLR	(SEQ	ID	NO:	214)
LRPVFKNTSVGPLYSGCRLTLLR	(SEQ	ID	NO:	215)
LRSLFKSTSVGPLYSGCRLTLLR	(SEQ	ID	NO:	216)
LRSLFKSTSVGPLYSGCRLTSLR	(SEQ	ID	NO:	217)
LTPLFKNTSVGPLYSGCRLTLLR	(SEQ	ID	NO:	218)
LTPLFRNTSVSSLYSGCRLTLLR	(SEQ	ID	NO:	219)
LMPLFKNTSVSSLYSGCRLTLLR	(SEQ	ID	NO:	220)
RPLFOKSSM.GPFYLGCOLTSLR	(SEO	ID	NO:	221)

### Exon 3

66 123				
${\tt PEKDSSAMAVDAICTHRPDPEDLGLDRERLYWELSNLTNGIQELGPYTLDRNSLYVNG}$	(SEQ	ID	NO:	222)
${\tt PEKDGAATGVDAICTHRLDPKSPGLNREQLYWELSKLTNDIEELGPYTLDRNSLYVNG}$	(SEQ	ID	NO:	223)
${\tt PKKDGAATGVDAICTHRLDPKSPGLNREQLYWELSKLTNDIEELGPYTLDRNSLYVNG}$	(SEQ	ID	NO:	224)
${\tt PEKDGTATGVDAICTHHPDPKSPRLDREQLYWELSQLTHNITELGHYALDNDSLFVNG}$	(SEQ	ID	NO:	225)
${\tt PEKDGEATGVDAICTHRPDPTGPGLDREQLYLELSQLTHSITELGPYTLDRDSLYVNG}$	(SEQ	ID	NO:	226)
${\tt PEKDGAATGMDAVCLYHPNPKRPGLDREQLYWELSQLTHNITELGPYSLDRDSLYVNG}$	(SEQ	ID	NO:	227)
${\tt PEKDGAATGMDAVCLYHPNPKRPGLDREQLYCELSQLTHNITELGPYSLDRDSLYVNG}$	(SEQ	ID	NO:	228)
${\tt PEKDGAATRVDAACTYRPDPKSPGLDREQLYWELSQLTHSITELGPYTLDRVSLYVNG}$	(SEQ	ID	NO:	229')
${\tt PKKDGAATKVDAICTYRPDPKSPGLDREQLYWELSQLTHSITELGPYTQDRDSLYVNG}$	(SEQ	ID	NO:	230)
${\tt PKKDGAATKVDAICTYRPDPKSPGLDREQLYWELSQLTHSITELGPYTQDRDSLYNVG}$	(SEQ	ID	NO:	231)
${\tt PEKDGAATRVDAVCTHRPDPKSPGLDRERLYWKLSQLTHGITELGPYTLDRHSLYVNG}$	(SEQ	ID	NO:	232)
${\tt PEKDGVATRVDAICTHRPDPKIPGLDRQQLYWELSQLTHSITELGPYTLDRDSLYVNG}$	(SEQ	ID	NO:	233)
${\tt SEKDGAATGVDAICIHHLDPKSPGLNRERLYWELSQLTNGIKELGPYTLDRNSLYVNG}$	(SEQ	ID	NO:	234)
${\tt SEKDGAATGVDAICTHRLDPKSPGLDREQLYWELSQLTNGIKELGPYTLDRNSLYVNG}$	(SEQ	ID	NO:	235)
${\tt SEKDGAATGVDAICTHRLDPKSPGVDREQLYWELSQLTNGIKELGPYTLDRNSLYVNG}$	(SEQ	ID	NO:	236)
${\tt SEKDGAATGVDAICTHRVDPKSPGVDREQLYWELSQLTNGIKELGPYTLDRNSLYVNG}$	(SEQ	ID	NO:	237)
${\tt SEKDGAATGVDAICTHHLNPQSPGLDREQLYWQLSQMTNGIKELGPYTLDRNSLYVNG}$	(SEQ	ID	NO:	238)
${\tt PEKRGAATGVDTICTHRLDPLNPGLDREQLYWELSKLTRGIIELGPYLLDRGSLYVNG}$	(SEQ	ID	NO:	239)
${\tt PEKNGAATGMDAICSHRLDPKSPGLNREQLYWELSQLTHGIKELGPYTLDRNSLYVNG}$	(SEQ	ID	NO:	240)
${\tt PEKNGAATGMDAICSHRLDPKSPGLDREQLYWELSQLTHGIKELGPYTLDRNSLYVNG}$	(SEQ	ID	NO:	241)
${\tt PEKHGAATGVDAICTLRLDPTGPGLDRERLYWELSQLTNSVTELGPYTLDRDSLYVNG}$	(SEQ	ID	NO:	242)
${\tt PEKHGAATGVDAICTLRLDPTGPGLDRERLYWELSQLTNSITELGPYTLDRDSLYVNG}$	(SEQ	ID	NO:	243)
${\tt PEKHEAATGVDTICTHRVDPIGPGLDRERLYWELSQLTNSITELGPYTLDRDSLYVNG}$	(SEQ	ID	NO:	244)
${\tt PEKQEAATGVDTICTHRVDPIGPGLDRERLYWELSQLTNSITELGPYTLDRDSLYVNG}$	(SEQ	ID	NO:	245)
${\tt PEKQEAATGVDTICTHRVDPIGPGLDRERLYWELSQLTNSITELGPYTLDRDSLYVDG}$	(SEQ	ID	NO:	246)
${\tt PEKDKAATRVDAICTHHPDPQSPGLNREQLYWELSQLTHGITELGPYTLDRDSLYVDG}$	(SEQ	ID	NO:	247)
${\tt SVKNGAETRVDLLCTYLQPLSGPGLPIKQVFHELSQQTHGITRLGPYSLDKDSLYLNG}$	(SEQ	ID	NO:	248)
${\tt PEKDGAATGVDTTCTYHPDPVGPGLDIQQLYWELSQLTHGVTQLGFYVLDRDSLFING}$	(SEQ	ID	NO:	249)

Figure 7C

Exon 4					Exon 5					
124 135					136	156				
FTHRSSMPTTST	(SEQ	ID	NO:	250)	PGTSTVDVGTSGTPSSSI	PSPT	(SEQ	ID	NO:	278)
FTHRSSMPTTSI	(SEQ	ID	NO:	251)	PGTSTVDLRTSGTPSSLS	SSPTIM	(SEQ	ID	NO:	279)
FTHRTSVPTSST	(SEQ	ID	NO:	252)	PGTSTVDLGTSGTPFSLI	PSPA	(SEQ	ID	NO:	280)
FTHRTSVPTTST	(SEQ	ID	NO:	253)	PGTSTVDLG.SGTPSSLI	PSPT	(SEQ	ID	NO:	281)
FTHRSSVPTTSS	(SEQ	ID	NO:	254)	PGTSTVDLG.SGTPSLPS	SSPT	(SEQ	ID	NO:	282)
FTHRSSVSTTST	(SEQ	ID	NO:	255)	PGTSTVDLGTSGTPSSLI	PSPT	(SEQ	ID	NO:	283)
FTHRSSVAPTST	(SEQ	ID	NO:	256)	PGTPTVDLGTSGTPVSK	PGPS	(SEQ	ID	NO:	284)
FTHRSSGLTTST	(SEQ	ID	NO:	257)	PWTSTVDLGTSGTPSPVF	PSPT	(SEQ	ID	NO:	285)
FTHRSFGLTTST	(SEQ	ID	NO:	258)	PGTSTVYWATTGTPSSF	GHT	(SEQ	ID	NO:	286)
FTHRSSFLTTST	(SEQ	ID	NO:	259)	PGTSTVHLATSGTPSSLE	GHT	(SEQ	ID	NO:	287)
FTHRNFVPITST	(SEQ	ID	NO:	260)	PGTSTVHLATSGTPSPLE	GHT	(SEQ	ID	NO:	288)
FTHRSSVPTTSI	(SEQ	ID	NO:	261)	PDTSTMHLATSRTPASLS	GPT	(SEQ	ID	NO:	289)
FTHQSSVSTTST	(SEQ	ID	NO:	262)	PGTSAVHLETSGTPASLE	PGHT	(SEQ	ID	NO:	290)
FTHQTSAPNTST	(SEQ	ID	NO:	263)	PGTSAVHLETTGTPSSFE	GHT	(SEQ	ID	NO:	291)
FTHQTFAPNTST	(SEQ	ID	NO:	264)	PGTSTVHLGTSETPSSLF	PRPI	(SEQ	ID	NO:	292)
FTHQNSVPTTST	(SEQ	ID	NO:	265)	PGTSIVNLGTSGIPPSLF	ETT	(SEQ	ID	NO:	293)
FTHQSSMTTTRT	(SEQ	ID	NO:	266)	PGTFTVQPETSETPSSLF	GPT	(SEQ	ID	NO:	294)
FTHWIPVPTSST	(SEQ	ID	NO:	267)	PGTPTVDLGTSGTPVSKP	GPS	(SEQ	ID	NO:	295)
FTHWSPIPTTST	(SEQ	ID	NO:	268)	PGTPTVYLGASKTPASIF	GPS	(SEQ	ID	NO:	296)
FTHWSSGLTTST	(SEQ	ID	NO:	269)	PKPATTFLPPLSEATT		(SEQ	ID	NO:	297)
FHPRSSVPTTST	(SEQ	ID	NO:	270)	QINFHIVNWNLSNPDPTS	SEY	(SEQ	ID	NO:	298)
${\tt FNPRSSVPTTST}$	(SEQ	ID	NO:	271)						
FNPWSSVPTTST	(SEQ	ID	NO:	272)						
FTQRSSVPTTSI	(SEQ	ID	NO:	273)						
FTQRSSVPTTST	(SEQ	ID	NO:	274)						
FTQRSSVPTTSV	(SEQ	ID	NO:	275)						
YNEPGLDEPPTT	(SEQ	ID	NO:	276)						
YAPQNLSIRGEY	(SEQ	ID	NO:	277)						



901 951 1001 1051 1101 1201 1201 1301 1401 1451 1451 1601 1651	ISATEPÄVPE SPHESEATAS WYHPAVTST TVPRTTPNYS HSEPDTTPSI	ARSPGAEATS DEPTITUSED VEDMVASQVA SSGTDTSITI PALALSSGEE	EPTTSFIÎVS ETHÎSSAIPÎ LPVSPGASKM LTSLVIŜSGÎ DSÎTÎFPÎLÎ	EPPYEPETTA IQLIHPAETN PMVPRFPKF SHSKSDTTLP VAITSPGPEA	SSAVSTÄTIS PDMSDLVTSL VPSSGÄDTST ÄFPÄLSEÄPY EPETÄATWLÄ	HPAETSTIVS GIIPNESHRG SDIAPSMVPS PGVDIRSGVP ITIIPPSIPG	VVÄSQVÄSSA TDÄSTAIPÄL ÄPSPGEPETÄ ASSATHPGTQ TGFTVPIRÄV	PSSEPDTMAS WVTHPPQTST PVSRTTSSFS HSSPDATPVM ATSPRTEASS	AVLTÄI SPGA PEMÖTSQIÄS SGAAÄSTIVP TIÄHSPGMPE TTALLSÄHPR	FETSKTFFPAS PVFPQVSETT ASLFIRPGAE ISTALPTQTT SSLFTLLVTG	tsrvdlspfa SpgvSaktap LSfhpgfets fmipfSflsl gllettglla	FSSAETSTS TLTLTVSPAV SGLSSASITÄ DKPQTVTSWN FETSPSVTSV	GPPEFSRIVI GITMLIPSE MPPPFKISHG EGVSPTILR ITMVEATNLA	PPGSSPFVAK ITITENTLAG SLFTPLTFPG MSTLASESVT SRISYNHRSW	ISTISSYNRR YWTPATSTPV TSTFSPGIST SSIPSSTAAT VPFMVPFTLN	FTITNLQYEE DWRHPGSRKF NATERELQGL LKPLFRNSSL EYLYSGCRLA	SLRPEKDSSA MAVDAICTHR PDPEDLGLDR ERLYWELSNL INGIQELGPY
					-												
EHITKIPNE AAHRGTIRPV KGPQTSTSPA SPKGLHTGGT  TTTTTALKTT SLATLTTSVY TPTLGTLTPL NASRQMASTI FPDVPETTS SLATSLGAET STALPRTPS VLNRESETTA PVIQTLDVS SSEPDTTASW VIHPAETIPT VSKTTPPFT SHGADVSA IPTNISPSEL DALTPLVTIS GTDTSTTFPT TTTTLTHPAE TSSTIPRTIP NESHHESDAT BSIATSPGAE SPGAEDLVTS QVTSSGTDRN MIPTLTLSP GEPKTIASLV PPTSTISPAV SRLVTSMVTS LAAKTSTTNR ALTNSPGEPA TSPTISPAV SRLVTSMVTS LAAKTSTTNR ALTNSPGEPA TSPTISPAV STIPLITLS PGEPETTPSM ATSHGEEASS TPGVVTSLVT STALPTLTLSSGEP TPPSMATSH TLEVPGMVT SLVASSRAVT STILPTLTLTLS PGEPETTPSM TSPGVVTSLVT SSRAVTSTT PILTFSLGEP TTPSMATSH TLEVPGWVT SLVASSRAVT STILPTLTLTLS TASSAVPTPT TASSAVPTPT TASSAVPTPT TASSAVPTPT TASSAVPTPT TASSAVTSTT TASSAVTST TASSAVTS T																	
EHITKIPNE AAHRGTIRPV I TTTTALKTT SRATLITESVY TEPDVPETTS SLATSLGAET SPOTOTLOVS SSEPDTTASW SHGADVSA IPTNISPSEL I TTWLTHPAE TSSTIPRTIP I PREABLUTS QUESGTDRN I POSTISPAV SRLVTSKOT STEPHSKSDT STOTSSRAVT STIPLITES I PREVPGWYT SIPRISPSEL I PREVPGWYT SIPRISPSEL I PREVPGWYT SIPRISPSEL I PREVPGWYT SIPRISPSEL I PREVPGWYT SIVASSRAVT STIPLITES I PREVPGWYT SIVASSRAVT STASSRAVTSTI I PREVPGWYT SIVASSRAVT STASSRAVPTPT SIPRISPSEL I PREVPGWYT SIVASSRAVT SIPRISPSEL I PREVPGWYT SIVASSRAVT SIPRISPSEL I PREVPGWYT SIVASSRAVT SIPRISPSEL I PREVPGWYT SIPRISPSEL I PREVPGENSSI ANTONSPERV OO O O O O O O O O O O O O O O O O O	KGPQTSTSPA SPKGLHTGGT	PPLGTLTPL NASROMASTI	STALPRITES VINRESETTA	VIHPAETIPT VSKTTPNFFH	OALTPLVTIS GTDTSTTFPT	NFSHHESDAT PSIATSPGAE	MILTLILSP GEPKTIASLV	LAAKTSTTNR ALTNSPGEPA	PSMTTSHGA ESSSAVPTPT	PGEPETTPSM ATSHGEEASS	PILTFSLGEP ETTPSMATSH	STILPILIS PGEPETTPSM	SGVNSTSIPT LILSPGELET	PLVTSSRAVT STTIPILTLS	GMVTSLVTSS RAVISTTIPT	VSPTVQGLVT SLVTSSGSET	VPTLTVSTGE PFTNISLVTH
	>	, YV	SAET	TASW 1	PSEL 1	PRTIP !	GTDRN 1	TSMVTS ]	HSKSDT	PILTLS 1	VISTI	SSRAVT :	SLVTSS	VSGVVT	VSPEVP (	AIPTLA	TEASSV

# Stucture of Carboxy Terminal Domain

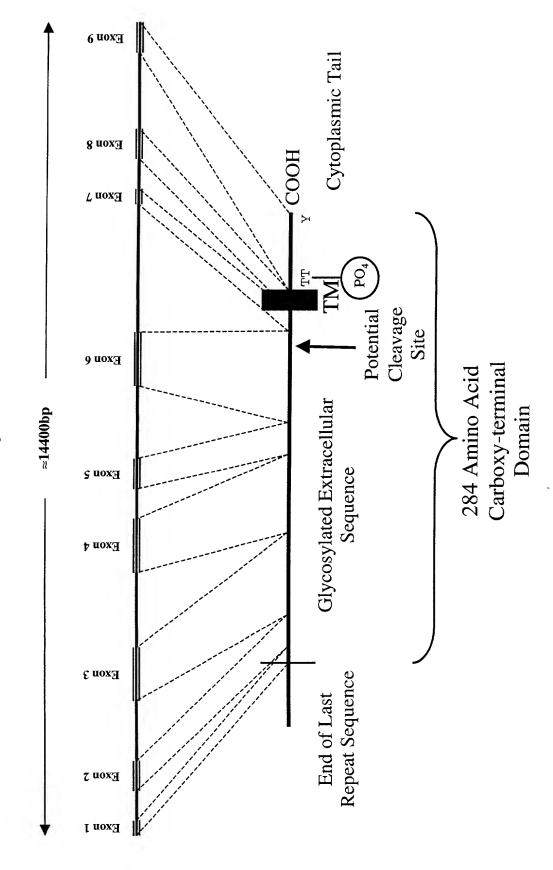


Figure 9A

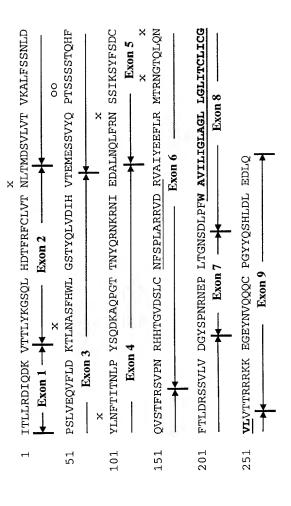


Figure 9B (SEQ ID NO: 300)

## Proposed Structure of CA125

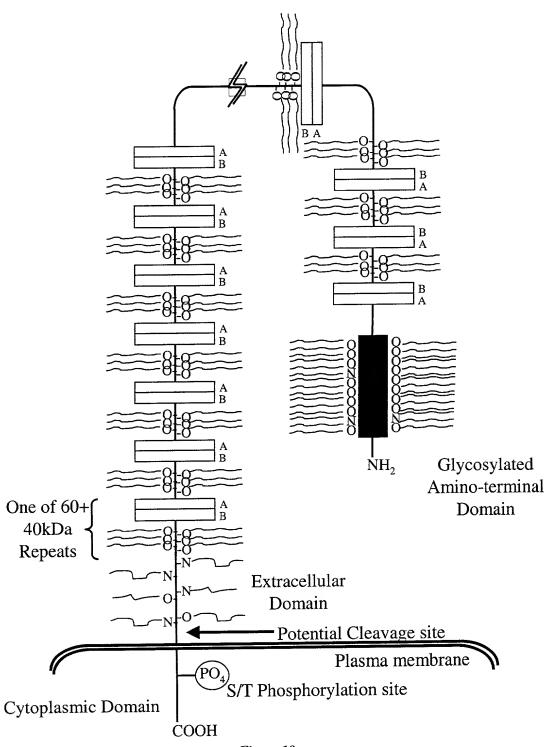


Figure 10